R. Leman

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RAW SEQUENCE LISTING

DATE: 11/28/2000 TTME: 15:11:49

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Input Set : A:\Intracel 58138-084.ST25.txt Output Set: N:\CRF3\11282000\1043530.raw

PATENT APPLICATION: US/09/043,530

DEC 15 2000.

TECH CENTER 1600/2900

ENTERED

```
2 <110> APPLICANT: Intracel Corporation
             Pilkington, Glenn
              Gilmour, Page
              Chanock, Robert
              Crowe, James
              Murphy, Brian
      9 <120> TITLE OF INVENTION: Neutralizing Monoclonal Antibodies to Respiratory Syncytial Virus
    11 <130> FILE REFERENCE: 58138-084
    13 <140> CURRENT APPLICATION NUMBER: US 09/043,530
C--> 15 <141> CURRENT FILING DATE: 2000-11-14
    17 <160> NUMBER OF SEQ ID NOS: 24
    19 <170> SOFTWARE: PatentIn version 3.0
    21 <210> SEQ ID NO: 1
    23 <211> LENGTH: 123
    25 <212> TYPE: PRT
    27 <213> ORGANISM: Homo sapiens
    29 <220> FEATURE:
    31 <221> NAME/KEY: domain
    33 <222> LOCATION: (1)..(123)
    35 <400> SEQUENCE: 1
    37 Leu Glu Glu Ser Gly Gly Asp Leu Val Gln Pro Gly Arg Ser Leu Arg
                                            10
    40~{\rm Leu~Ser~Cys~Ser~Thr~Ser~Gly~Phe~Ser~Phe~Gly~Asp~Tyr~Pro~Val~Asn}
    41
                   20
                                        25
    43 Trp Phe Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Leu Gly Ile Val
              35
    46 Arg Ser Arg Leu Tyr Gly Gly Thr Leu Gln Tyr Ala Ala Ser Val Glu
         50
                               5.5
    49 Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Ser Ile Ala Tyr Leu
                            70
    52 His Met Asn Ser Leu Lys Ser Glu Asp Thr Ala Val Tyr Tyr Cys Gly
                                            90
                       8.5
    55 Val Pro Val Ala Asn Ile Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr
                   100
                                       1.05
    58 Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Ser
              1.15
                                    1.20
    61 <210> SEQ ID NO: 2
    63 <211> LENGTH: 27
    65 <212> TYPE: PRT
    67 <213> ORGANISM: Homo sapiens
    69 <220> FEATURE:
W--> 71 <221> NAME/KEY: region
    73 <222> LOCATION: (1)..(27)
    75 <400> SEQUENCE: 2
    77 Leu Glu Glu Ser Gly Gly Asp Leu Val Gln Pro Gly Arg Ser Leu Arg
    78 1.
                       5
                                            1.0
    80 Leu Ser Cys Ser Thr Ser Gly Phe Ser Phe Gly
```



Input Set : A:\Intracel 58138-084.ST25.txt
Output Set: N:\CRF3\11282000\I043530.raw

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83 <210> SEQ ID NO: 3
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    87 <212> TYPE: PRT
     89 <213> ORGANISM: Homo sapiens
     91 <220> FEATURE:
W--> 93 <221> NAME/KEY: region
    95 <222> LOCATION: (1)..(5)
    97 <400> SEQUENCE: 3
     99 Asp Tyr Pro Val Asn
    100 1
    102 <210> SEQ ID NO: 4
    104 <211> LENGTH: 14
    106 <212> TYPE: PRT
    108 <213> ORGANISM: Homo sapiens
    110 <220> FEATURE:
W--> 112 <221> NAME/KEY: region
    114 <222> LOCATION: (1)..(14)
    116 <400> SEQUENCE: 4
    118 Trp Phe Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Leu Gly
    119 1
    121 <210> SEQ ID NO: 5
    123 <211> LENGTH: 19
    125 <212> TYPE: PRT
127 <213> ORGANTSM: Homo sapiens
    129 <220> FEATURE:
W--> 131 <221> NAME/KEY: region
    133 <222> LOCATION: (1)..(19)
    135 <400> SEQUENCE: 5
    137 Ile Val Arg Ser Arg Leu Tyr Gly Gly Thr Leu Gln Tyr Ala Ala Ser
    140 Val Glu Gly
    143 <210> SEQ ID NO: 6
    145 <211> LENGTH: 32
    147 <212> TYPE: PRT
    149 <213> ORGANISM: Homo sapiens
    151 <220> FEATURE:
W--> 153 <221> NAME/KEY: region
    155 <222> LOCATION: (1)..(32)
    157 <400> SEQUENCE: 6
    159 Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Ser Ile Ala Tyr Leu His
    160 1 5
                                        1.0
                                                              15
    162 Met Asn Ser Leu Lys Ser Glu Asp Thr Ala Val Tyr Tyr Cys Gly Val
    1.63 20
    165 <210> SEQ ID NO: 7
    167 <211> LENGTH: 7
    169 <212> TYPE: PRT
    171 <213> ORGANISM: Homo sapiens
    173 <220> FEATURE:
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Input Set : A:\Intracel 58138-084.ST25.txt
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    177 <222> LOCATION: (1)..(7)
    179 <400> SEQUENCE: 7
    181 Pro Val Ala Asn Ile Asp Tyr
    182 1
                      5
    184 <210> SEQ ID NO: 8
    186 <211> LENGTH: 19
    188 <212> TYPE: PRT
    190 <213> ORGANTSM: Homo sapiens
    192 <220> FEATURE:
W--> 194 <221> NAME/KEY: region
    196 <222> LOCATION: (1)..(19)
    198 <400> SEQUENCE: 8
    200 Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly
             5
    203 Pro Ser Ser
    206 <210> SEQ ID NO: 9
    208 <211> LENGTH: 109
    210 <212> TYPE: PRT
    212 <213> ORGANISM: Homo sapiens
    214 <220> FEATURE:
    216 <221> NAME/KEY: domain
    218 <222> LOCATION: (1)..(109)
    220 <400> SEQUENCE: 9
    222 Leu Thr Gln Pro His Ser Val Ser Glu Ser Leu Gly Lys Thr Val Thr
    223 1
                   5
                               10
    225 Ile Ser Cys Thr Arg Ala Gly Gly Ser Ile Ala Ser Asn Tyr Val Gln
    226 20 25
    228 Trp Tyr Gln Gln Arg Pro Gly Ser Ser Pro Thr Thr Val Ile Tyr Glu
    229 35
                                40
    231 Asp Asn Gin Arg Pro Phe Gly Val Pro Asp Arg Phe Ser Gly Ser Ile
                            55
    234 Asp Thr Ser Ser Asm Ser Ala Ser Leu Thr Ile Ser Gly Leu Lys Thr
    235 65
                       7.0
                                  75
    237 Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Ser Glu Asn Pro
                  8.5
                                  90
    240 Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly 241 100 105
    241
                100
                                   105
    243 <210> SEQ ID NO: 10
    245 <211> LENGTH: 19
    247 <212> TYPE: PRT
    249 <213> ORGANISM: Homo sapiens
    251 <220> FEATURE:
W--> 253 <221> NAME/KEY: region
    255 <222> LOCATION: (1)..(19)
    257 <400> SEQUENCE: 1.0
    259 Leu Thr Gln Pro His Ser Val Ser Glu Ser Leu Gly Lys Thr Val Thr
    260 1
                    5
                                        10
    262 Ile Ser Cys
```

Input Set : A:\Intracel 58138-084.ST25.txt
Output Set: N:\CRF3\11282000\1043530.raw

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265 <210> SEQ ID NO: 11
     267 <211> LENGTH: 13
     269 <212> TYPE: PRT
     271 <213> ORGANISM: Homo sapiens
     273 <220> FEATURE:
W--> 275 <221> NAME/KEY: region
     277 <222> LOCATION: (1)..(13)
     279 <400> SEQUENCE: 11
     281 Thr Arg Ala Gly Gly Ser Ile Ala Ser Asn Tyr Val Gln
     284 <210> SEQ ID NO: 12
     286 <211> LENGTH: 15
     288 <212> TYPE: PRT
     290 <213> ORGANISM: Homo sapiens
     292 <220> FEATURE:
W--> 294 <221> NAME/KEY: region
     296 <222> LOCATION: (1)..(15)
     298 <400> SEQUENCE: 12
     300 Trp Tyr Gln Gln Arg Pro Gly Ser Ser Pro Thr Thr Val Ile Tyr
                                  10
     301 1
                       5
     303 <210> SEQ ID NO: 13
     305 <211> LENGTH: 7
     307 <212> TYPE: PRT
     309 <213> ORGANISM: Homo sapiens
     311 <220> FEATURE:
W--> 313 <221> NAME/KEY: region
     315 <222> LOCATION: (1)..(7)
     317 <400> SEQUENCE: 13
     319 Glu Asp Asn Gln Arg Pro Phe
     320 1
     322 <210> SEQ TD NO: 14
     324 <21.1> LENGTH: 34
     326 <212> TYPE: PRT
     328 <213> ORGANISM: Homo sapiens
     330 <220> FEATURE:
W--> 332 <221> NAME/KEY: region
    334 <222> LOCATION: (1)..(14)
336 <400> SEQUENCE: 14
     338 Gly Val Pro Asp Arg Phe Ser Gly Ser Ile Asp Thr Ser Ser Asn Ser
                        5
                                            10
    341 Ala Ser Leu Thr Ile Ser Gly Leu Lys Thr Glu Asp Glu Ala Asp Tyr 342 20 25 30
     344 Tyr Cys
     347 <210> SEQ ID NO: 15
     349 <211> LENGTH: 10
     351 <212> TYPE: PRT
     353 <213> ORGANISM: Homo sapiens
    355 <220> FEATURE:
W--> 357 <221> NAME/KEY: region
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Input Set : A:\Intracel 58138-084.ST25.txt
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359 <222> LOCATION: (1)..(10)
     361 <400> SEQUENCE: 15
     363 Gln Ser Tyr Asp Ser Glu Asn Pro Trp Val
     364 1
     366 <210> SEQ ID NO: 16
     368 <211> LENGTH: 11
     370 <212> TYPE: PRT
     372 <213> ORGANISM: Homo sapiens
     374 <220> FEATURE:
W--> 376 <221> NAME/KEY: region
     378 <222> LOCATION: (1)..(11)
     380 <400> SEQUENCE: 16
     382 Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly
     383 1.
     385 <210> SEQ ID NO: 17
     387 <211> TENGTH: 17
389 <212> TYPE: DNA
     391 <213> ORGANISM: synthetic primer
     393 <220> FEATURE:
     395 <221> NAME/KEY: primer_bind
     397 <222> LOCATION: (1)..(17)
     399 <400> SEQUENCE: 17
     400 attaaccctc actaaag
                                                                                     17
     403 <210> SEO ID NO: 18
405 <211> LENGTH: 21
     407 <212> TYPE: DNA
     409 <213> ORGANISM: synthetic primer
     411 <220> FEATURE:
     413 <221> NAME/KEY: primer_bind
     415 <222> LOCATION: (1)..(21)
     417 <400> SEQUENCE: 18
     418 gaattotaaa etagotagto g
                                                                                     21
     421 <210> SEQ ID NO: 19
     423 <211> LENGTH: 18
     425 <212> TYPE: DNA
     427 <213> ORGANISM: synthetic primer
    429 <220> FEATURE:
431 <221> NAME/KEY: primer_bind
     433 <222> LOCATION: (1)..(18)
     435 <400> SEQUENCE: 19
     436 gaagtagtee ttgaccag
                                                                                     1.8
    439 <210> SEQ ID NO: 20
441 <211> LENGTH: 21
     443 <212> TYPE: DNA
     445 <213> ORGANISM: synthetic primer
     447 <220> FEATURE:
     449 <221> NAME/KEY: primer_bind
     451 <222> LOCATION: (1)..(21)
     453 <400> SEQUENCE: 20
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/043,530

DATE: 11/28/2000

TIME: 15:11:50

Input Set : A:\Intracel 58138-084.ST25.txt
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L:71 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:93 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:112 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:131 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
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